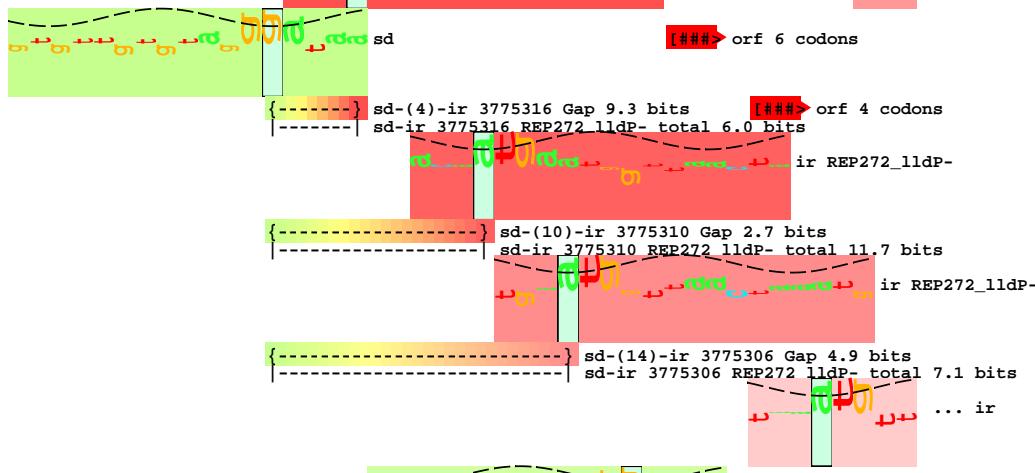


- 1 -

piece 1, NC_000913, REP272_11dP-, config: linear, direction: -, begin: 3775451, end: 3775101

Sequence alignment diagram showing the amino acid sequence of a protein across different organisms. The top row shows the gene sequence with start sites (*), stop sites (asterisks), and frame shifts (red dots). The bottom row shows the deduced amino acid sequence for each organism, with identical residues grouped by color.



p35 2.7 bits [###> orf 6 codons

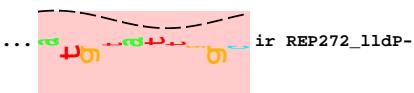
{-----} sd-(6)-ir 3775249 Gap 4.3 bits

sd-ir 3775249 REP272_lldP- total 5.5 bits

{-----} p35-(24)-p10 3775263 Gap 2-4 bits

- 2 -

|-----| p35-p10 3775263 total 4.8 bits



```
... } sd-(14)-ir 3775289 Gap 4.9 bits  
... | sd-ir 3775289 REP272_11dP- total 5.4 bits
```